

# Heterogeneity of natural Foxp3<sup>+</sup> T cells: A committed regulatory T-cell lineage and an uncommitted minor population retaining plasticity

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**Natural regulatory T cells (T<sub>reg</sub>) represent a distinct lineage of T lymphocytes committed to suppressive functions, and expression of the transcription factor Foxp3 is thought to identify this lineage specifically. Here we report that, whereas the majority of natural CD4<sup>+</sup>Foxp3<sup>+</sup> T cells maintain stable Foxp3 expression after adoptive transfer to lymphopenic or lymphoreplete recipients, a minor fraction enriched within the CD25<sup>+</sup> subset actually lose it. Some of those Foxp3<sup>+</sup> T cells adopt effector helper T cell (T<sub>h</sub>) functions, whereas some retain "memory" of previous Foxp3 expression, reacquiring Foxp3 upon activation. This minority "unstable" population exhibits flexible responses to cytokine signals, relying on transforming growth factor- $\beta$  to maintain Foxp3 expression and responding to other cytokines by differentiating into effector T<sub>h</sub> in vitro. In contrast, CD4<sup>+</sup>Foxp3<sup>+</sup>CD25<sup>high</sup> T cells are resistant to such conversion to effector T<sub>h</sub> even after many rounds of cell division. These results demonstrate that natural Foxp3<sup>+</sup> T cells are a heterogeneous population consisting of a committed T<sub>reg</sub> lineage and an uncommitted subpopulation with developmental plasticity.**

immunological tolerance | lineage commitment | transcription factor

Compelling evidence indicates that natural T<sub>reg</sub>, initially identified as CD4<sup>+</sup>CD25<sup>+</sup> T cells, are central to the establishment and maintenance of immunological self-tolerance and immune homeostasis (1). Such cells express predominantly, if not specifically, the transcription factor Foxp3 (2–4). Studies in mice engineered to express a Foxp3 reporter confirmed that Foxp3 expression is confined to a subset of  $\alpha\beta$  T cells and correlates with suppressor activity irrespective of CD25 expression (5, 6). Furthermore, mutations of the Foxp3 gene result in the development of a catastrophic lymphoproliferative autoimmune disorder in mice and humans due to defective generation of functional T<sub>reg</sub> (3, 4), whereas ectopic Foxp3 expression in conventional CD4<sup>+</sup> T cells confers a phenotype and function similar to, if not identical to, natural T<sub>reg</sub> (2–4). More recently, continuous and high levels Foxp3 expression was found necessary for maintenance of T<sub>reg</sub> phenotype and function (7, 8). These findings collectively led to the notion that Foxp3 represents a specific and faithful molecular marker to identify the T<sub>reg</sub> lineage and acts as its "specification factor" or "master regulator" (2, 5).

Early studies of autoimmunity provoked by neonatal thymectomy in mice (1), and of transplantation tolerance to tissue grafts induced by previous transplantation of pure thymic epithelium in birds and in mice (9), indicated that natural T<sub>reg</sub> represent a distinct thymus-committed lineage. Although it has also become evident that further Foxp3<sup>+</sup> T<sub>reg</sub> can be generated from peripheral naïve CD4<sup>+</sup> T cells upon "tolerogenic" antigen presentation in vivo or activation in the presence of transforming growth factor (TGF)- $\beta$  *in vitro* (10, 11), this notion of a stable lineage has been strengthened by data showing that the phenotypic and functional features of CD4<sup>+</sup>CD25<sup>+</sup> T<sub>reg</sub> are stably maintained after many rounds of cell

division in vitro and in vivo (1). In line with this, the majority, if not all, of CD4<sup>+</sup>CD25<sup>+</sup> T cells appear to maintain Foxp3 expression when transferred into normal lymphoreplete mice, and their Foxp3 stability has been associated with chromatin remodeling of the Foxp3 locus (12). All these studies examined phenotypic and functional stability of CD4<sup>+</sup>CD25<sup>+</sup> T cells at the population level, leaving it unclear as to whether absolutely all natural Foxp3<sup>+</sup> T cells were so fixed in their behavior.

In contrast, others have claimed that Foxp3 expression may not be entirely specific for the T<sub>reg</sub> lineage and that natural T<sub>reg</sub> may have a much more plastic phenotype. First, human naïve CD4<sup>+</sup> T cells were shown to up-regulate FOXP3 transiently at low levels after in vitro TCR stimulation without acquiring T<sub>reg</sub> characteristics (13). Second, Foxp3 expression in murine TGF- $\beta$ -induced Foxp3<sup>+</sup> T cells (inducible T<sub>reg</sub> or iT<sub>reg</sub>) has been claimed to be unstable and to be readily lost upon secondary stimulation (12). Third, some natural Foxp3<sup>+</sup> T cells from Foxp3 reporter mice were recently shown to down-regulate Foxp3 and "transdifferentiate" into interleukin (IL)-17-producing effector T<sub>h</sub> under the influence of IL-6 and autocrine TGF- $\beta$  (14, 15). These observations clearly challenge the prevailing notion, yet the root of this apparent contradiction is unclear.

This study was undertaken to evaluate the stability of Foxp3 expression and T<sub>reg</sub> phenotype of CD4<sup>+</sup>Foxp3<sup>+</sup> T cells isolated from nonmanipulated normal mice in vivo and in vitro. We demonstrate heterogeneity within natural Foxp3<sup>+</sup> T cells and provide evidence that they comprise a majority committed T<sub>reg</sub> lineage and a minor subpopulation with developmental plasticity.

## Results

**Some Peripheral CD4<sup>+</sup>Foxp3<sup>+</sup> T Cells Lose Foxp3 Expression Under Lymphopenic and Lymphoreplete Conditions.** CD4<sup>+</sup>EGFP<sup>+</sup> and EGFP<sup>−</sup> T cells were sorted from the spleen and LN of Foxp3<sup>EGFP</sup> mice and transferred into RAG2<sup>−/−</sup> mice either alone or mixed at a 1:1 or 1:10 ratio. To distinguish the two donor populations in the co-transferred group, EGFP<sup>+</sup> and EGFP<sup>−</sup> cells were obtained from Foxp3<sup>EGFP</sup> Ly5.2 and Ly5.1 congenic mice, respectively. Four weeks after transfer, LN (Fig. 1A) and spleen (not depicted) were analyzed for EGFP expression in donor CD4<sup>+</sup> T cells. Approximately 50% of donor CD4<sup>+</sup> T cells were EGFP<sup>−</sup> in the recipients of EGFP<sup>+</sup> cells alone. The generation of EGFP<sup>−</sup> cells from EGFP<sup>+</sup>

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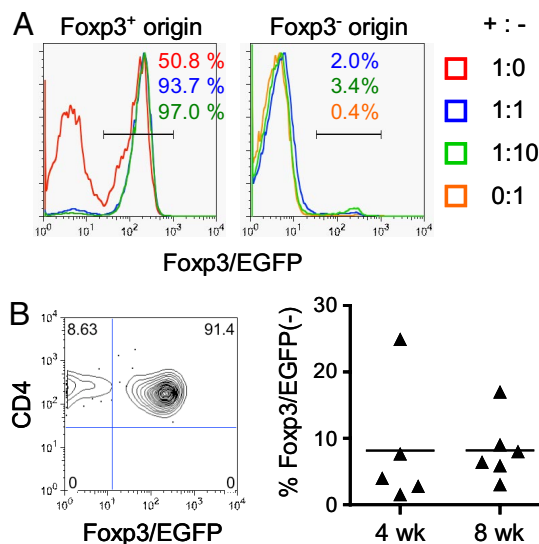
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**Fig. 1.** A fraction of  $CD4^{+}Foxp3^{+}$  T cells lose Foxp3 expression in lymphopenic and normal lymphoreplete conditions. (A)  $CD4^{+}EGFP^{+}$  and  $EGFP^{-}$  T cells were sorted from  $Foxp3^{EGFP}$  Ly5.2 and  $Foxp3^{EGFP}$  Ly5.1 mice, respectively, and adoptively transferred into  $RAG2^{-/-}$  mice, either alone ( $1 \times 10^5$  each) or mixed at a 1:1 or 1:10 ratio ( $1 \times 10^5$   $EGFP^{+}$  plus  $1 \times 10^5$  or  $1 \times 10^6$   $EGFP^{-}$ ). Four weeks after transfer, LN cells were stained for CD4, TCR $\beta$  and Ly5.1. Shown are representative EGFP expression profiles of the  $Foxp3^{+}$  (Ly5.1 $^{+}$ ) or  $Foxp3^{-}$  (Ly5.1 $^{+}$ ) donor-derived  $CD4^{+}TCR\beta^{+}$  cells recovered from the indicated host mice. (B)  $1 \times 10^5$   $CD4^{+}EGFP^{+}$  T cells sorted from  $Foxp3^{EGFP}$  Ly5.1/Thy1.2 mice were adoptively transferred into Ly5.2/Thy1.1 host mice 4 and 8 weeks earlier. Pooled LN and spleen cells were first enriched for donor T cells by depleting Thy1.1 $^{+}$ , Ig $^{+}$ , and adherent cells by panning, and were stained for Ly5.1, Thy1.2, and CD4. A representative EGFP and CD4 profile of the Ly5.1 $^{+}$ Thy1.2 $^{+}$  donor cells (left) and frequencies of  $EGFP^{-}$  cells in  $CD4^{+}Ly5.1^{+}Thy1.2^{+}$  donor T cells (right) are shown. Each symbol indicates individual host mouse.

cells was not caused by  $EGFP^{-}$  contaminants in the inocula but reflected Foxp3 down-regulation, as small numbers of Ly5.1  $EGFP^{-}$  T cells that had been mixed into the sorted Ly5.2  $EGFP^{+}$  T cells (manipulated such that similar numbers of Ly5.1 and Ly5.2  $EGFP^{-}$  cells were injected) failed to “grow out” within the recipients [supporting information (SI) Fig. S1]. When co-transferred with  $EGFP^{-}$  cells, the numbers of  $EGFP^{+}$  donor cells that maintained EGFP expression (hereafter denoted as  $Foxp3^{+}$  T cells) increased in a manner depending on the numbers of co-injected  $EGFP^{-}$  cells (Fig. 1A, Fig. S2A), indicating that  $Foxp3^{-}$  cells promoted expansion and/or survival of  $Foxp3^{+}$  cells. In contrast, the numbers of  $EGFP^{+}$  donor cells that had lost EGFP expression ( $Foxp3^{+}$  T cells) were reduced when co-transferred with  $EGFP^{-}$  T cells, indicating that  $Foxp3^{-}$  cells inhibited the expansion/survival of  $Foxp3^{+}$  T cells or the down-regulation of Foxp3 expression. The loss of Foxp3 expression and its inhibition by  $Foxp3^{-}$  T cells was also observed when  $CD3e^{-/-}$  mice were used as hosts, indicating that the presence of B cells does not affect Foxp3 down-regulation (Fig. S2B). As higher numbers of donor T cells could be obtained from  $CD3e^{-/-}$  hosts compared with  $RAG2^{-/-}$  hosts,  $CD3e^{-/-}$  mice were used in some of the following analyses when large numbers of donor-derived T cells were needed.

To determine whether Foxp3 down-regulation from  $CD4^{+}Foxp3^{+}$  T cells takes place under lymphoreplete conditions, we adoptively transferred  $CD4^{+}EGFP^{+}$  T cells sorted from  $Foxp3^{EGFP}$  Ly5.1/Thy1.2 mice into Ly5.2/Thy1.1 mice. Use of the dual congenic markers as well as pre-enrichment of donor cells allowed us to faithfully detect small numbers of donor T cells as Ly5.1 $^{+}$ Thy1.2 $^{+}$  cells. Approximately 8% of  $EGFP^{+}$  donor T cells were  $EGFP^{-}$  when analyzed at 4 and 8 weeks after transfer (Fig. 1B).

These results demonstrate that some  $Foxp3^{+}$  T cells had down-

regulated Foxp3 expression in both lymphopenic and lymphoreplete hosts, suggesting that Foxp3 down-regulation represents a normal physiological process. Inasmuch as the frequencies of  $Foxp3^{+}$  T cells were low in the presence of  $Foxp3^{-}$  “competitor” T cells, they must normally be a somewhat rare population.

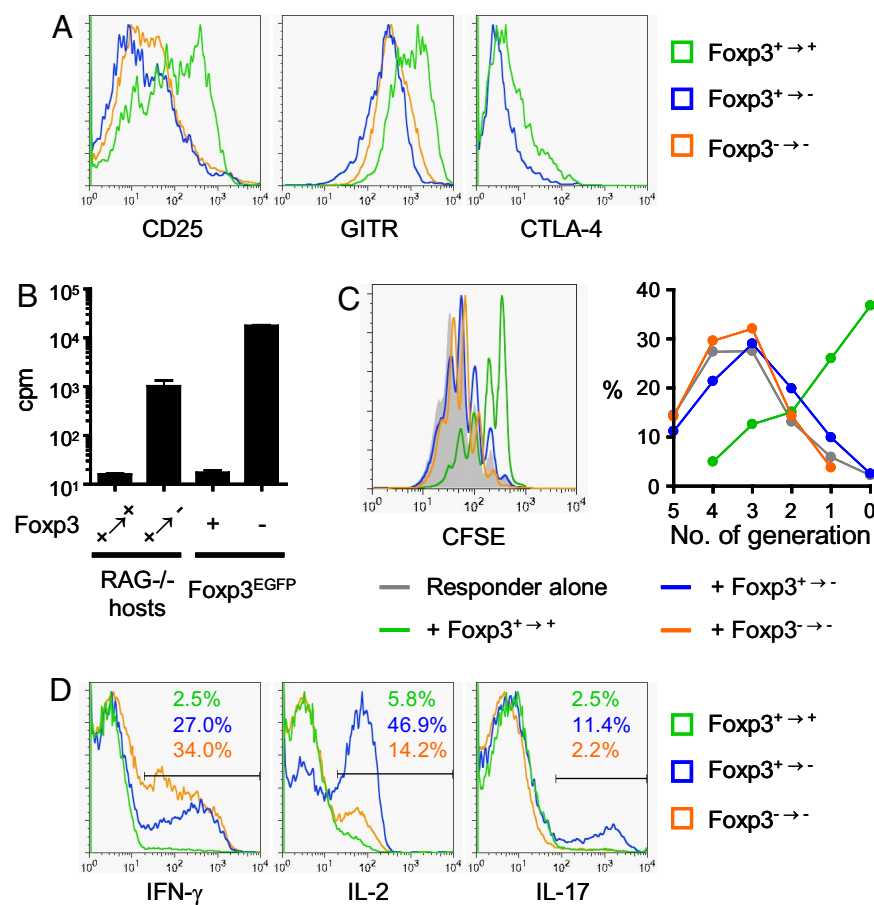
**The Majority of Foxp3-Down-Regulated T Cells Lose  $T_{reg}$  Phenotype and Function and Produce Inflammatory Cytokines.** We next examined whether  $CD4^{+}Foxp3^{+}$  T cells lose  $T_{reg}$  phenotype and function upon loss of Foxp3 expression and, if so, whether they differentiate into effector  $T_h$ . CD25, GITR, and CTLA-4, well-known  $T_{reg}$  phenotypic markers were less abundant on  $Foxp3^{+}$  T cells than on  $Foxp3^{+}$  T cells, and comparable to control  $Foxp3^{-}$  T cells recovered from  $RAG2^{-/-}$  recipients of  $Foxp3^{+}$  T cells alone ( $Foxp3^{+}$  T cells) (Fig. 2A). When stimulated in vitro with anti-CD3, both  $Foxp3^{+}$  T cells as well as  $EGFP^{+}$  T cells from  $Foxp3^{EGFP}$  mice failed to proliferate, indicating, as expected, that  $Foxp3^{+}$  T cells maintained their anergic state (Fig. 2B). In contrast,  $Foxp3^{+}$  T cells proliferated to a greater extent than  $Foxp3^{+}$  T cells, yet much less than  $EGFP^{-}$  T cells from  $Foxp3^{EGFP}$  mice, indicating a partial anergy (Fig. 2B). When co-cultured with  $CD4^{+}CD25^{-}$  responder T cells,  $Foxp3^{+}$  T cells did not suppress their proliferation as efficiently as  $Foxp3^{+}$  T cells (Fig. 2C). However, responder T cells proliferated slightly less in the presence of  $Foxp3^{+}$  T cells than in their absence, whereas  $Foxp3^{+}$  T cells enhanced responder T-cell proliferation. These results indicate that  $Foxp3^{+}$  T cells had lost their suppressive activity, although not completely.

To examine whether  $Foxp3^{+}$  T cells had differentiated into effector  $T_h$  in lymphopenic hosts,  $Foxp3^{+}$ ,  $Foxp3^{+}$ , and  $Foxp3^{+}$  T cells were stimulated ex vivo and analyzed for their cytokine production at the single cell level (Fig. 2D). Many  $Foxp3^{+}$  T cells produced significant IL-2, IL-17, and IFN- $\gamma$  compared with  $Foxp3^{+}$  T cells. Notably,  $Foxp3^{+}$  T cells preferentially produced IL-2 or IL-17 compared with  $Foxp3^{+}$  T cells, which mainly produced IFN- $\gamma$ . None of these populations produced IL-4 at a significant frequency (<1%) (not depicted). These results suggest that at least some  $Foxp3^{+}$  T cells had differentiated into effector  $T_h$ .

**A Significant Fraction of Foxp3-Down-Regulated T cells Reacquire Foxp3 Expression in Vitro and in Vivo.** In the above in vitro suppression assay, we noted that a fraction of  $Foxp3^{+}$  T cells re-expressed EGFP, whereas  $Foxp3^{+}$  T cells or  $Foxp3^{-}$  T cells from  $Foxp3^{EGFP}$  mice did not (not shown). This preferential Foxp3 re-induction from  $Foxp3^{+}$  T cells was also seen when stimulated with anti-CD3/CD28 mAbs-coated beads (Fig. 3A).  $Foxp3^{+}$  T cells re-expressed Foxp3 in a TGF- $\beta$ -independent manner, as neutralizing concentrations of anti-TGF- $\beta$  mAbs and TGF $\beta$ RII-Fc chimeric proteins did not affect the outcome (Fig. 3B). In contrast, TCR stimulation was essential, because  $Foxp3^{+}$  T cells failed to reacquire EGFP expression without anti-CD3/CD28 beads (not shown). The observed re-induction of Foxp3 expression in a fraction of  $Foxp3^{+}$  T cells may well explain their partially anergic phenotype and weak suppressor activity (Fig. 2B and C) despite a large number of IL-2-producing cells (Fig. 2D).

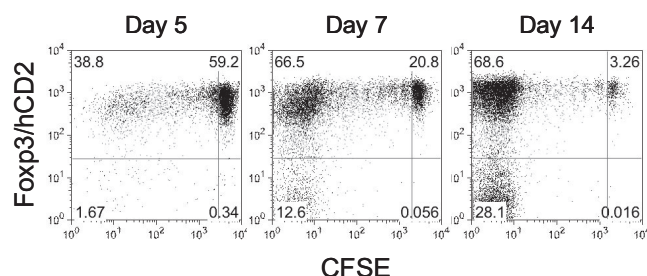
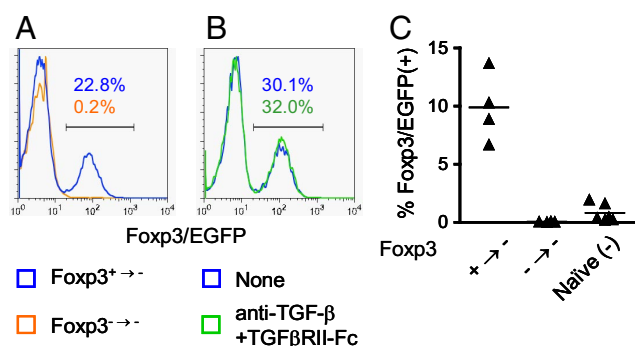
To address whether  $Foxp3^{+}$  T cells preferentially reacquire Foxp3 expression in vivo, we injected  $Foxp3^{+}$  T cells or control  $Foxp3^{+}$  T cells from  $CD3e^{-/-}$  hosts into secondary  $CD3e^{-/-}$  hosts. We found that  $\approx 10\%$  of  $Foxp3^{+}$  T cells re-expressed EGFP, whereas few  $Foxp3^{+}$  T cells and only  $\approx 1\%$  of naïve  $EGFP^{-}$  T cells up-regulated EGFP (Fig. 3C).

These results indicate that  $Foxp3^{+}$  T cells are heterogeneous and comprise not only effector  $T_h$  but also putative  $T_{reg}$  precursors poised for Foxp3 transcription.

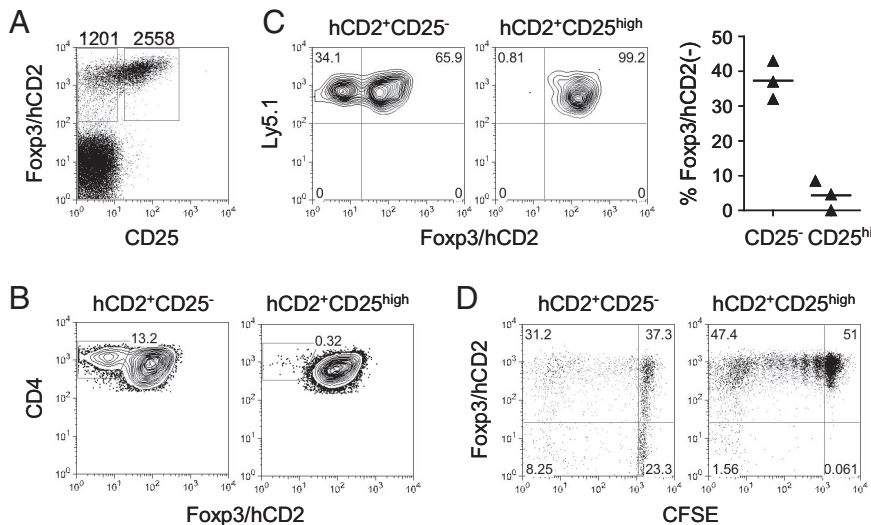


**Foxp3-Down-Regulated T Cells Are Generated by Extensive Lymphopenia-Driven Expansion of a Rare Subpopulation.** We observed that Foxp3<sup>+</sup>→<sup>-</sup> T cells constituted ≈50% of Foxp3<sup>+</sup> donor-derived cells in lymphopenic mice, whereas they represented a minor population when co-transferred with Foxp3<sup>-</sup> T cells or when transferred into normal mice. We wondered whether the “unsta-

ble” subpopulation was better able to expand in lymphopenic conditions when competitor  $\text{Foxp3}^-$  T cells were absent. We therefore examined the relationship between  $\text{Foxp3}$  downregulation and cell division using a  $\text{Foxp3}^{\text{hCD2}}$  reporter that we have established recently. Sorted  $\text{CD4}^+\text{hCD2}^+$  T cells were labeled with CFSE, injected into  $\text{RAG2}^{-/-}$  mice, and analyzed for hCD2 expression simultaneously with CFSE dilution on days 5, 7, and 14 after transfer. On day 5, only  $\approx 2\%$  of donor cells were  $\text{hCD2}^-$ ; but  $\approx 83\%$  of this small number of  $\text{Foxp3}^{+\rightarrow-}$  T cells had already divided more than once whereas  $\approx 60\%$  of  $\text{Foxp3}^{+\rightarrow+}$  T cells still remained undivided, indicating enrichment of proliferating cells in  $\text{Foxp3}^{+\rightarrow-}$  T cells (Fig. 4). The frequencies of  $\text{hCD2}^-$  T cells progressively increased through days 7 to 14, and all of the







**Fig. 5.** CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>-</sup> T cells exhibit lower and less stable Fopx3 expression than CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>high</sup> T cells. (A) LN cells from Fopx3<sup>hCD2</sup> mice were stained for hCD2, CD4, and CD25. Representative CD25<sup>-</sup> and hCD2 expression profiles on CD4<sup>+</sup> cells are shown. The number above each gate indicates the mean fluorescent intensity of hCD2. (B) Sorted CD4<sup>+</sup>hCD2<sup>+</sup>CD25<sup>-</sup> and CD25<sup>high</sup> T cells were stimulated in vitro with anti-CD3/CD28 beads in the presence of IL-2 and analyzed for hCD2 and CD4 expression on day 3. (C)  $1 \times 10^5$  Ly5.1 CD4<sup>+</sup>hCD2<sup>+</sup>CD25<sup>-</sup> and CD25<sup>high</sup> T cells were transferred into RAG2<sup>-/-</sup> mice, and LN cells were stained for hCD2, Ly5.1, and Ly5.2 on day 5. Shown are representative hCD2 and Ly5.1 expression profiles (left) and frequencies of hCD2<sup>-</sup> cells (right) in donor (Ly5.1<sup>+</sup>Ly5.2<sup>-</sup>) cells. (D) Ly5.1 CD4<sup>+</sup>hCD2<sup>+</sup>CD25<sup>-</sup> and CD25<sup>high</sup> T cells were stained with CFSE and transferred into Fopx3<sup>hCD2</sup> Ly5.2 mice ( $4 \times 10^5$  cells). On day 14, pooled LN and spleen cells were stained for Ly5.1, Ly5.2, and hCD2 and enriched for donor Ly5.1<sup>+</sup> cells by magnetic sorting before analysis. Representative CFSE and hCD2 profiles on Ly5.1<sup>+</sup>Ly5.2<sup>-</sup> cells are shown.

Fopx3<sup>+</sup> T cells had lost CFSE, indicating that they had undergone multiple rounds of cell division. These results suggest that the Fopx3<sup>+</sup> T cells represent progeny of a relatively rare subpopulation of Fopx3<sup>+</sup> T cells that undergo extensive expansion in lymphopenic conditions. However, it remains possible that any Fopx3<sup>+</sup> T cells can lose Fopx3 expression with a certain probability, requiring a defined number of divisions to lose Fopx3.

To distinguish these two possibilities, we compared the TCR repertoires of Fopx3<sup>+</sup> and Fopx3<sup>+</sup> T cells by flow cytometry using mAbs for V $\alpha$ 2 and 6 V $\beta$  chains. If only a rare subpopulation of Fopx3<sup>+</sup> T cells lost Fopx3 and then expanded, their TCR repertoires would be more oligoclonal than Fopx3<sup>+</sup> T cells and the two repertoires should largely be distinct. In contrast, if any Fopx3<sup>+</sup> T cells can down-regulate Fopx3 after a defined number of cell divisions, the two repertoires ought to be similar to each other. As shown in Fig. S3, although the patterns of V $\alpha$ /V $\beta$  usage in polyclonal Fopx3<sup>+</sup> T cells were stable among different individual mice, those in both Fopx3<sup>+</sup> T cells and Fopx3<sup>+</sup> T cells were highly variable, indicative of oligoclonal expansion. However, the variation in Fopx3<sup>+</sup> T cells was significantly larger than in Fopx3<sup>+</sup> T cells, suggesting the former to be more oligoclonal than the latter. Moreover, within each individual mouse, Fopx3<sup>+</sup> T cells and Fopx3<sup>+</sup> T cells showed distinct V $\alpha$ /V $\beta$  usage patterns, suggesting that clones expanding in the two populations were different. These results support the notion that Fopx3<sup>+</sup> T cells represent progeny of a relatively rare subpopulation of Fopx3<sup>+</sup> T cells.

**CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>-</sup> T Cells Exhibit Unstable, Whereas CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>high</sup> T Cells Exhibit Stable, Fopx3 Expression.** Let us suppose that Fopx3<sup>+</sup> T cells in the normal T-cell repertoire harbor a minor population of unstable Fopx3<sup>+</sup> T cells. How might these be distinguished from stable Fopx3<sup>+</sup> T cells? We asked whether Fopx3 stability could be associated with the expression of a Fopx3-dependent marker. We found that CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>-</sup> T cells expressed lower levels of the hCD2 reporter than did CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>high</sup> T cells (Fig. 5A). The former subset also expressed GITR and CTLA-4 at lower levels than the latter (not shown). In line with their lower Fopx3 expression, CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>-</sup> T cells were less anergic and suppressive than CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>high</sup> T cells (Fig. S4). These results led us to hypothesize that CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>-</sup> T cells might be the ones harboring uncommitted T<sub>reg</sub>, the Fopx3 expression of which is labile.

We sorted CD4<sup>+</sup>Fopx3<sup>+</sup> T cells into CD25<sup>-</sup> and CD25<sup>high</sup> cells and tested their Fopx3 stability in vitro and in vivo. Fopx3<sup>hCD2</sup> mice were used instead of Fopx3<sup>EGFP</sup> mice in these analyses, because the

hCD2 reporter enables better separation of Fopx3<sup>low</sup> cells from Fopx3<sup>+</sup> cells due to its high sensitivity. In addition, pre-enrichment of hCD2<sup>+</sup> cells by magnetic sorting followed by FACS sorting yielded a highly pure (>99%) population of this small subset. When hCD2<sup>+</sup>CD25<sup>-</sup> and CD25<sup>high</sup> T cells were stimulated in vitro with anti-CD3/CD28 beads in the presence of IL-2, we found that  $\approx 11\%$  of the former population lost hCD2 expression whereas >99% of the latter maintained hCD2 (Fig. 5B). Furthermore, when transferred into RAG2<sup>-/-</sup> mice,  $\approx 37\%$  of Fopx3<sup>+</sup>CD25<sup>-</sup> T cells preferentially down-regulated hCD2, whereas most of Fopx3<sup>+</sup>CD25<sup>high</sup> T cells maintained hCD2 (Fig. 5C).

When labeled with CFSE and transferred into Fopx3<sup>hCD2</sup> Ly5.2 mice, the vast majority of hCD2<sup>+</sup>CD25<sup>high</sup> T cells maintained hCD2 expression even after multiple cycles of cell division, although 1–2% of them down-regulated hCD2 while losing CFSE (Fig. 5D). In contrast,  $\approx 32\%$  of hCD2<sup>+</sup>CD25<sup>-</sup> T cells lost hCD2 expression, the majority ( $\approx 75\%$ ) of which did so without cell division. This lack of cell division of Fopx3<sup>+</sup> T cells accounts for their low representation in nonlymphopenic conditions and in turn supports the idea that they are generated by extensive proliferation of a rare subset of Fopx3<sup>+</sup> T cells in lymphopenic conditions.

These results collectively indicate that the unstable cells come from a rare subpopulation enriched within the CD25<sup>-</sup> subset of Fopx3<sup>+</sup> T cells.

**CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>-</sup> T Cells Contain Uncommitted T<sub>reg</sub> That Retain T<sub>h</sub> Potential, Whereas CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>high</sup> T Cells Represent a Committed T<sub>reg</sub> Lineage.** Our results suggest that some of CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>-</sup> T cells remain uncommitted to T<sub>reg</sub> fate and retain options on alternative T<sub>h</sub> pathways, whereas CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>high</sup> T cells represent a committed T<sub>reg</sub> lineage. Developmental plasticity of uncommitted progenitors can be revealed when they are placed in environments that are permissive for alternative fate choices. It is well established that cytokines provide such “environments.” TGF- $\beta$  is known to induce Fopx3 expression and to instruct “iT<sub>reg</sub>” differentiation, whereas IL-12, IL-4, and IL-6 together with TGF- $\beta$  promote T<sub>h</sub>1, T<sub>h</sub>2, and T<sub>h</sub>17 differentiation, respectively. We asked whether CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>-</sup> and CD25<sup>high</sup> T cells depended on TGF- $\beta$  for their Fopx3 maintenance and whether the above cytokines would encourage differentiation into effector T<sub>h</sub> subsets.

CD4<sup>+</sup>Fopx3<sup>+</sup>CD25<sup>-</sup> and CD25<sup>high</sup> T cells were stimulated with anti-CD3/CD28 beads in the presence or absence of TGF- $\beta$ 1 or anti-TGF- $\beta$  mAbs (Fig. 6A). The proportion of Fopx3<sup>+</sup>CD25<sup>-</sup> T cells that lost hCD2 expression increased 3-fold with stimulation in the presence of anti-TGF- $\beta$ , whereas they not only maintained but also up-regulated hCD2 in the presence of TGF- $\beta$ 1. In contrast, few



T cells from Foxp3 transgenic mice lose Foxp3 expression and differentiate into effector  $T_h$  when activated under polarizing conditions in vitro (16), although it was not clear whether they were derived from Foxp3<sup>+</sup> T cells or small numbers of Foxp3<sup>+</sup> T cells that were present in those mice. We have extended these observations to natural Foxp3<sup>+</sup> T cells from normal mice and demonstrated that they have the potential to differentiate into effector  $T_h$  including  $T_h17$  (14, 15). On the other hand, however, our results are incompatible with the view that natural Foxp3<sup>+</sup> T cells represent a fully plastic population in that any cell can be converted to effector  $T_h$ . Instead, the majority of natural Foxp3<sup>+</sup> T cells, particularly the CD25<sup>+</sup> subset, exhibited stable Foxp3 expression even after extensive proliferation and were resistant to conversion into effector  $T_h$ , so reflecting a committed  $T_{reg}$  lineage that has lost alternative  $T_h$  options.

We found that Foxp3-down-regulated T cells preferentially re-acquire Foxp3 expression upon in vitro activation or secondary transfer, indicating that they retain “memory” of previous Foxp3 expression. This Foxp3 re-induction was TGF- $\beta$ -independent, suggesting that its underlying mechanism is distinct from that of *de novo* Foxp3 induction in naïve T cells. Chromatin modifications including selective demethylation of CpG motifs have been implicated in epigenetic imprinting of gene expression, as shown for cytokine genes (17). Indeed, CD4<sup>+</sup>CD25<sup>+</sup> T cells were recently shown to display such modifications at the Foxp3 locus, particularly complete demethylation of one of the enhancers (12). Importantly, TGF- $\beta$ -induced Foxp3<sup>+</sup> T cells and activated human T cells, both of which show unstable Foxp3 expression, displayed little demethylation, suggesting that the demethylated Foxp3 locus represents a specific signature of the committed  $T_{reg}$  lineage (12, 18). In light of the present study, we can speculate that these T cells with “Foxp3 memory” may belong to the committed  $T_{reg}$  lineage, in which their Foxp3 locus has been remodeled and now “accessible”, yet transiently being inactive for transcription. Because TCR stimulation was required for Foxp3 re-induction, we suggest that TCR signals may keep Foxp3 transcription active within such a remodeled locus.

Finally, the findings of this study have important implications with respect to the role of Foxp3 in  $T_{reg}$  lineage commitment. Since some Foxp3<sup>+</sup> T cells can be converted to effector  $T_h$ , Foxp3 alone is clearly insufficient (although necessary) to commit precursor cells to the  $T_{reg}$  lineage. Our data thus support the emerging view that the  $T_{reg}$  lineage is determined by a higher-order regulation operating upstream of, and in cooperation with, Foxp3 (19, 20). However, the molecular nature of such a regulatory system is yet to be uncovered.

## Materials and Methods

**Mice.** C57BL/6J (B6) mice were obtained from Clea Japan. B6.Foxp3<sup>EGFP</sup> (21), B6.Foxp3<sup>hCD2</sup>, B6.RAG2<sup>-/-</sup>, B6.CD3 $\epsilon$ <sup>-/-</sup>, B6.Thy1.1, and B6.Ly5.1 congenic mice were bred under specific pathogen-free conditions in our animal facility and used at 5–12 weeks of age. B6.Foxp3<sup>hCD2</sup> reporter mice were generated by homologous recombination in a B6-derived ES cell line using a targeting construct in which cDNA encoding a human CD2-human CD52 fusion protein along with an intra-ribosomal entry site was inserted into the 3' untranslated region of the endogenous Foxp3 locus. Details of the generation and characterization of Foxp3<sup>hCD2</sup> mice will be reported elsewhere. All animal experiments were performed in accordance with approved protocols from the Institutional Animal Care at RIKEN.

**Reagents.** Purified and conjugated monoclonal antibodies (mAbs) were purchased from BD Bioscience, eBioscience, or R&D Systems, and recombinant proteins were obtained from R&D Systems or Peprotech. (See *SI Materials and Methods* for details.)

**Lymphocyte Preparations, Flow Cytometry, Cell Sorting.** Single-cell suspensions from spleen and LN were prepared as described previously (22). Flow-cytometric analyses were performed as described (22) using a FACSCalibur instrument (Becton Dickinson). Cell sorting was performed using a FACSaria cell sorter (Becton Dickinson). (See *SI Materials and Methods* for details.) The purity of the sorted populations was invariably >99%.

**T-Cell Functional Assays.** For proliferation assays, T cells ( $2 \times 10^4$ /well) were stimulated with 0.5  $\mu$ g/ml anti-CD3 in the presence of T cell-depleted,  $\gamma$ -irradiated B6 spleen cells as antigen-presenting cells ( $8 \times 10^4$ /well) for 3 days as described (2). Results are shown as mean [<sup>3</sup>H]thymidine incorporation  $\pm$  SD in triplicate cultures. For suppression assays, CFSE-labeled responder T cells were stimulated alone or together with  $T_{reg}$  ( $5 \times 10^4$  each/well) for 3 days with 0.5  $\mu$ g/ml anti-CD3 in the presence of antigen-presenting cells ( $2 \times 10^5$ /well) in 96-well U-bottom plates. To examine Foxp3 stability and  $T_h$  differentiation, T cells ( $1 \times 10^5$ /well) were stimulated with anti-CD3/anti-CD28 mAbs-coated beads (Dynal) at a 1:1 cell:bead ratio in 96-well, flat-bottom plates. Where indicated, IL-2 (10 ng/ml), IL-4 (10 ng/ml), IL-6 (20 ng/ml), IL-12 (10 ng/ml), TGF- $\beta$ 1 (5 ng/ml), anti-TGF- $\beta$ 1/2/3 (30  $\mu$ g/ml), and/or TGF $\beta$ R1I-Fc (3  $\mu$ g/ml) were added. To assess cytokine production, cells were stimulated for 4 hours with 50 ng/ml PMA and 500 ng/ml ionomycin in the presence of brefeldin A and then fixed, permeabilized, and stained with anti-cytokine or isotype control mAbs as described elsewhere (2).

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